

| Project Title   | Funding     | Strategic Plan Objective | Institution                                    |
|---|-------------|--------------------------|--|
| MSSNG   | \$5,802,895 | Q3.L.B                   | Autism Speaks (AS)                             |
| Simons Simplex Collection support grant   | \$5,983     | Q3.L.B                   | Baylor College of Medicine                     |
| Accelerating Autism Genetics via Whole Population Ascertainment in Denmark                        | \$0         | Q3.L.B                   | Broad Institute, Inc.                          |
| Pieces of the Puzzle: Uncovering the Genetics of Autism   | \$1,699,790 | Q3.L.B                   | Broad Institute, Inc.                          |
| 2/4-The Autism Sequencing Consortium: Autism gene discovery in >20,000 exomes                     | \$157,618   | Q3.S.A                   | BROAD INSTITUTE, INC.                          |
| 2/4-The Autism Sequencing Consortium: Autism gene discovery in >20,000 exomes                     | \$415,893   | Q3.S.A                   | BROAD INSTITUTE, INC.                          |
| Use of High-throughput Splicing Assays to Prioritize Autism Gene Candidates                       | \$62,500    | Q3.L.B                   | Brown University                               |
| Interactome perturbation by large-scale mutagenesis to find risk variants - Project 2             | \$29,831    | Q3.Other                 | Carnegie Mellon University                     |
| An exploration of genetic and behavioral variables in Autism Spectrum Disorder                    | \$18,200    | Q3.S.A                   | Center for Autism and Related Disorders (CARD) |
| Autism genetics: homozygosity mapping and functional validation                                   | \$765,736   | Q3.L.B                   | CHILDREN'S HOSPITAL CORPORATION                |
| Discovery and Functional Characterization of Gene Regulatory Networks (GRNs) of Autism Risk Genes | \$59,900    | Q3.Other                 | CHILDREN'S HOSPITAL OF LOS ANGELES             |
| The Future of Genomics Medicine in Patient Care: Contributions from CHOP                          | \$906,296   | Q3.L.B                   | Children's Hospital of Philadelphia            |
| Genetic basis of autism   | \$4,000,000 | Q3.L.B                   | Cold Spring Harbor Laboratory                  |
| Elucidating pathogenic mutations disrupting RNA regulation in autism                              | \$225,000   | Q3.L.B                   | Columbia University                            |
| Identification and analysis of functional networks perturbed in autism                            | \$250,000   | Q3.L.B                   | Columbia University                            |
| Applications of novel statistical methods to CNVs in autism and schizophrenia                     | \$200,000   | Q3.L.B                   | Columbia University                            |
| Novel Statistical methods for DNA Sequencing Data, and applications to Autism.                    | \$318,575   | Q3.L.B                   | Columbia University                            |
| Interactome perturbation by large-scale mutagenesis to find risk variants ñ Core                  | \$97,702    | Q3.Other                 | Cornell University                             |
| FUNCTIONAL DISSECTION OF CNVS IN NEURODEVELOPMENTAL TRAITS  | \$366,666   | Q3.L.B                   | Duke University                                |
| Simons Simplex Collection support grant   | \$8,800     | Q3.L.B                   | Emory University                               |
| Autism Spectrum Disorders: Genomes to Outcomes  | \$0         | Q3.L.B                   | Hospital for Sick Children                     |
| Discovery of regulatory variants underlying pediatric neurological disease                        | \$0         | Q3.L.B                   | HudsonAlpha Institute for Biotechnology        |
| Integrating large scale whole exome data with whole genome data                                   | \$0         | Q3.L.B                   | ICAHN SCHOOL OF MEDICINE AT MOUNT SINAI        |
| Population-Based Autism Genetics & Environment Study  | \$165,663   | Q3.L.D                   | ICAHN SCHOOL OF MEDICINE AT MOUNT SINAI        |
| Population-Based Autism Genetics & Environment Study  | \$618,149   | Q3.L.D                   | ICAHN SCHOOL OF MEDICINE AT MOUNT SINAI        |
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| 1/4-The Autism Sequencing Consortium: Autism gene discovery in >20,000 exomes                                       | \$206,585   | Q3.S.A                   | ICAHN SCHOOL OF MEDICINE AT MOUNT SINAI   |
| 1/4-The Autism Sequencing Consortium: Autism gene discovery in >20,000 exomes                                       | \$720,372   | Q3.S.A                   | ICAHN SCHOOL OF MEDICINE AT MOUNT SINAI   |
| Illumina, Inc.  | \$0         | Q3.L.B                   | Illumina, Inc.  |
| Genomic profiling of autism families using whole-genome sequencing  | \$0         | Q3.L.B                   | Institut Pasteur  |
| Sequencing Female-enriched Multiplex Autism Families (FEMFs)  | \$0         | Q3.L.B                   | Johns Hopkins University  |
| Cryptic Genetic Causes of Autism  | \$141,719   | Q3.L.B                   | Massachusetts General Hospital  |
| Complex Genetic Architecture of Chromosomal Aberrations in Autism   | \$248,999   | Q3.L.B                   | Massachusetts General Hospital  |
| Role of the Intestinal Microbiome in Children with Autism   | \$0         | Q3.S.I                   | Massachusetts General Hospital  |
| Sequence-based discovery of genes with pleiotropic effects across diagnostic boundaries and throughout the lifespan | \$14,998    | Q3.L.B                   | Massachusetts General Hospital  |
| Simons Simplex Collection support grant   | \$10,000    | Q3.L.B                   | McGill University Health Centre- Montreal Children's Hospital                   |
| Genetic Epidemiology of Complex Traits  | \$747,204   | Q3.L.B                   | National Institutes of Health   |
| New York Genome Center, Inc.  | \$2,210,000 | Q3.L.B                   | New York Genome Center, Inc.  |
| Investigating the role of somatic mutations in autism spectrum disorders  | \$263,892   | Q3.L.B                   | OREGON HEALTH & SCIENCE UNIVERSITY  |
| Computational tools to analyze SNP data from patients with mental illness   | \$572,792   | Q3.L.B                   | PARTEK, INC.  |
| Identifying Patterns of Genetic Variants Conferring Risk for Neurodevelopmental Disorders                           | \$29,987    | Q3.L.B                   | Pennsylvania State University   |
| Whole-exome sequencing to identify causative genes for autism   | \$134,203   | Q3.L.B                   | ROCKEFELLER UNIVERSITY  |
| Advanced Autism Genetics: Biological Subgroups, Diagnostic Classification, and Resilience.                          | \$30,000    | Q3.S.A                   | State University of New York, Upstate Medical University                        |
| Thompson Center Clinical Site Network Pilot for the National Autism Cohort  | \$37,500    | Q3.L.B                   | The Curators of the University of Missouri                                      |
| High-throughput Screening of Novel Trinucleotide Repeat Expansion in Autism Spectrum Disorders                      | \$15,000    | Q3.L.B                   | The Hospital for Sick Children  |
| Extending ASD risk locus discovery to the non-coding genome - Core  | \$0         | Q3.L.B                   | The Regents of the University of California, San Francisco (Contracts & Grants) |
| Validation of candidate ASD genes by targeted sequencing with molecular inversion probes                            | \$101,258   | Q3.L.B                   | The Regents of the University of California, San Francisco (Contracts & Grants) |
| Exploring the Intersection of Autism and Homeostatic Synaptic Plasticity  | \$60,000    | Q3.Other                 | The Regents of the University of California, San Francisco (Contracts & Grants) |
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| Impact of Pten mutations on brain growth and social behavioral development.   | \$480,000   | Q3.S.K                   | The Scripps Research Institute                              |
| Extending ASD risk locus discovery to the non-coding genome - Project 1   | \$0         | Q3.L.B                   | The Trustees of Columbia University in the City of New York |
| Genomic influences on development and outcomes in infants at risk for autism  | \$0         | Q3.L.B                   | University of Alberta                                       |
| Genomic influences on developmental course and outcome in Infants at risk of ASD: A Baby Siblings Research Consortium (BSRC) Study  | \$0         | Q3.S.A                   | University of Alberta                                       |
| Genomic influences on development and outcomes in Infants at risk of ASD  | \$0         | Q3.S.A                   | University of Alberta                                       |
| A multi-platform approach to the functional assessment of ASD gene variants   | \$120,000   | Q3.Other                 | University of British Columbia                              |
| Simons Simplex Collection support grant   | \$13,200    | Q3.L.B                   | University of California, Los Angeles                       |
| Dosage effects of 22q11 region on autism-relevant neural systems  | \$0         | Q3.S.A                   | University of California, Los Angeles                       |
| Autism Genetics, Phase II: Increasing Representation of Human Diversity   | \$2,715,972 | Q3.S.D                   | University of California, Los Angeles                       |
| Rapid Phenotyping for Rare Variant Discovery in Autism  | \$453,878   | Q3.S.A                   | University of California, Los Angeles                       |
| Mutations in noncoding DNA and the missing heritability of autism   | \$244,030   | Q3.L.B                   | University of California, San Diego                         |
| Uncovering the Spectrum of De Novo Mutation in Autism through Whole Genome Sequencing   | \$35,000    | Q3.S.A                   | University of California, San Diego                         |
| 4/4 The Autism Sequencing Consortium: Autism gene discovery in >20,000 exomes   | \$676,656   | Q3.S.A                   | UNIVERSITY OF CALIFORNIA, SAN FRANCISCO                     |
| 4/4 The Autism Sequencing Consortium: Autism gene discovery in the >20,000 exomes (supplement)  | \$919,964   | Q3.S.A                   | UNIVERSITY OF CALIFORNIA, SAN FRANCISCO                     |
| Dosage effects of DUF1220 gene subtype CON1 in autism   | \$125,000   | Q3.L.B                   | University of Colorado, Denver                              |
| Simons Simplex Collection support grant   | \$9,159     | Q3.L.B                   | University of Illinois at Chicago                           |
| Determine risk genes enriched in intronic multispecies conserved sequences, and copy number variant breakpoints in autism overlap preferentially with short interspersed elements | \$21,000    | Q3.Other                 | University of Louisville                                    |
| Simons Simplex Collection support grant   | \$10,000    | Q3.L.B                   | University of Missouri                                      |
| University of North Carolina Clinical Site Network Pilot for the National Autism Cohort   | \$24,750    | Q3.L.B                   | University of North Carolina                                |
| Role of Selfish Spermatogonial Selection in Neurocognitive Disorders  | \$59,995    | Q3.L.B                   | University of Oxford  |
| 3/3-Sequencing Autism Spectrum Disorder Extended Pedigrees  | \$160,000   | Q3.L.B                   | University of Pennsylvania                                  |
| A history of behavioral genetics  | \$0         | Q3.Other                 | University of Pittsburgh                                    |

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| Interactome perturbation by large-scale mutagenesis to find risk variants - Project 1                  | \$24,172  | Q3.Other                 | University of Pittsburgh           |
| 3/4 - The Autism Sequencing Consortium: Autism gene discovery in >20,000 exomes                        | \$65,732  | Q3.S.A                   | University of Pittsburgh           |
| 3/4 - The Autism Sequencing Consortium: Autism gene discovery in >20,000 exomes                        | \$302,248 | Q3.S.A                   | University of Pittsburgh           |
| Combining WGS from Utah high-risk pedigrees and SSC families   | \$0       | Q3.L.B                   | University of Utah                 |
| 1/3 - Sequencing Autism Spectrum Disorder Extended Pedigrees   | \$298,000 | Q3.L.B                   | UNIVERSITY OF UTAH                 |
| Genome Sequencing pilot of Simons Simplex Collection   | \$0       | Q3.L.B                   | University of Washington           |
| Structural Variation and the Genetic Architecture of Autism  | \$0       | Q3.L.B                   | University of Washington           |
| Genetic basis of phenotypic variability in 16p11.2 deletion or duplication                             | \$285,856 | Q3.L.B                   | University of Washington           |
| University of Washington Clinical Site Network Pilot for the National Autism Cohort                    | \$37,500  | Q3.L.B                   | University of Washington           |
| Simons Simplex Collection support grant  | \$10,000  | Q3.L.B                   | University of Washington           |
| Autism subtypes by gene characterization   | \$318,824 | Q3.S.A                   | University of Washington           |
| Next Generation Gene Discovery in Familial Autism  | \$644,823 | Q3.L.B                   | University of Washington           |
| 2/3 Sequencing Autism Spectrum Disorder Extended Pedigrees   | \$231,750 | Q3.L.B                   | University of Washington           |
| The genetic basis underlying the phenotype heterogeneity of the 16p11.2 CNV                            | \$46,136  | Q3.S.A                   | University of Washington           |
| Sporadic Mutations and Autism Spectrum Disorders   | \$647,900 | Q3.S.A                   | University of Washington           |
| Simons Simplex Collection support grant  | \$8,912   | Q3.L.B                   | Vanderbilt University              |
| Genome-wide analysis of cis-regulatory elements in autism  | \$62,500  | Q3.L.B                   | Washington University in St. Louis |
| Quantifying Offspring ASD Risk for Unaffected Sisters of Males with ASD                                | \$35,000  | Q3.S.C                   | Washington University in St. Louis |
| Simons Simplex Collection support grant  | \$1,831   | Q3.L.B                   | Weill Cornell Medical College      |
| Simons Simplex Collection support grant  | \$10,000  | Q3.L.B                   | Yale University                    |
| Extending ASD risk locus discovery to the non-coding genome - Project 2                                | \$0       | Q3.L.B                   | Yale University                    |
| Integrating the genomics of Autism Spectrum Disorders(ASD) in consanguineous and "idiopathic" families | \$665,939 | Q3.L.B                   | Yale University                    |

